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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/714,449A

DATE: 10/02/2001

TIME: 09:23:49

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Input Set : A:\00431PHRM293.ST25.txt
Output Set: N:\CRF3\10022001\I714449A.raw

3 <110> APPLICANT: Vogeli, Gabriel
4 Huff, Rita
5 Sejlitz, Torsten
6 Lind, Peter
7 Slightom, Jerry
8 Schellin, Kathleen
9 Bannigan, Chris
10 Ruff, Valerie
11 Kaytes, Paul
12 Wood, Linda
13 Parodi, Luis
14 Hiebsch, Ronald
16 <120> TITLE OF INVENTION: Novel G Protein Coupled Receptors
18 <130> FILE REFERENCE: 00431PHRM293
20 <140> CURRENT APPLICATION NUMBER: 09/714,449A
21 <141> CURRENT FILING DATE: 2000-11-16
23 <150> PRIOR APPLICATION NUMBER: 60/165,838
24 <151> PRIOR FILING DATE: 1999-11-16
26 <150> PRIOR APPLICATION NUMBER: 60/198,568
27 <151> PRIOR FILING DATE: 2000-04-20
29 <150> PRIOR APPLICATION NUMBER: 60/166,071
30 <151> PRIOR FILING DATE: 1999-11-17
32 <150> PRIOR APPLICATION NUMBER: 60/166,678
33 <151> PRIOR FILING DATE: 1999-11-19
35 <150> PRIOR APPLICATION NUMBER: 60/173,396
36 <151> PRIOR FILING DATE: 1999-12-28
38 <150> PRIOR APPLICATION NUMBER: 60/184,129
39 <151> PRIOR FILING DATE: 2000-02-22
41 <150> PRIOR APPLICATION NUMBER: 60/185,421
42 <151> PRIOR FILING DATE: 2000-02-28
44 <150> PRIOR APPLICATION NUMBER: 60/185,554
45 <151> PRIOR FILING DATE: 2000-02-28
47 <150> PRIOR APPLICATION NUMBER: 60/186,530
48 <151> PRIOR FILING DATE: 2000-03-02
50 <150> PRIOR APPLICATION NUMBER: 60/186,811
51 <151> PRIOR FILING DATE: 2000-03-03
53 <150> PRIOR APPLICATION NUMBER: 60/188,114
54 <151> PRIOR FILING DATE: 2000-03-09
56 <150> PRIOR APPLICATION NUMBER: 60/190,310
57 <151> PRIOR FILING DATE: 2000-03-17
59 <150> PRIOR APPLICATION NUMBER: 60/190,800
60 <151> PRIOR FILING DATE: 2000-03-21
62 <150> PRIOR APPLICATION NUMBER: 60/201,190
63 <151> PRIOR FILING DATE: 2000-05-02
65 <150> PRIOR APPLICATION NUMBER: 60/203,111
66 <151> PRIOR FILING DATE: 2000-05-08
68 <150> PRIOR APPLICATION NUMBER: 60/207,094

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69 <151> PRIOR FILING DATE: 2000-05-25
 71 <160> NUMBER OF SEQ ID NOS: 190
 73 <170> SOFTWARE: PatentIn version 3.0
 75 <210> SEQ ID NO: 1
 76 <211> LENGTH: 1182
 77 <212> TYPE: DNA
 78 <213> ORGANISM: Homo sapiens

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 83 cctagctcct gctgatctag ttggggctcc agagtgggga ggagaaaggc actttgaaac 120
 85 ttctctgccc ttaccgtctt agccatcaaa ctctgagctg gagatagtga cgatgtgaca 180
 87 ggaactttcc ctgggcctct ctgggccaca attcctggcc gagagaaaga ggaggaatga 240
 89 ggtgagcacc ttcttctact ctaggggccat gtggttagagc tgcagtcgca cctccttctg 300
 91 ccaataggca tagatgagtg ggttgagcag ggagttgcc acgccgagca gccacaggta 360
 93 ccgttcacgc actaggtaga ggtgacactc ctggcaggcc acctgcacaa tgccagtgat 420
 95 aaggaagggg gtccaggata gagcaaagct cccaatgaga acagacacag tacggagagc 480
 97 tttgaagtgc ctgggagtcg gtggggatcg ataacctcca gccatggctc ctgcatgttc 540
 99 catctttcga atctgctggc tgtgcatgga ggcaatcttg agcatgtcgc agtagaagaa 600
 101 gacaaagagg agcatggctg ggaagaagcc aacgcaggag agggtcagca cgaagtgagg 660
 103 gtgaaatata gcaaagaagc tgcactgccc tttgtaggca gtctgctgga acatggggat 720
 105 tccgagtggg aggaagccaa tgaggtaga cactaaccac agcccggcaa tgcaggcccc 780
 107 ggccacgaac ccatcatga tcttcaagta gcggaagggc tgcttgatgg caaggtacct 840
 109 gtcaaaggtg atcagcatga ccgtgaggac agaggcagct gcggagggaag tgacaaatgc 900
 111 catccgcagg ctgcacaggg tcttctgtgt gggccgagaa gggctggaga gctggtctgt 960
 113 gagtaggcca gagatggcca caccaatcaa ggtgtcagcc acagccagat tcaaggtgaa 1020
 115 gcagagactg acaccatcat tcttgtggat caacagcagc acagccacag ccactagtgt 1080
 117 gttagtagca atgatgaggg aggccaggac agcaaggatc actccaaatg agaaagatga 1140
 119 ttccatgtct cgaagtggca ggacttcaat taccagggca tg 1182

122 <210> SEQ ID NO: 2

123 <211> LENGTH: 335

124 <212> TYPE: PRT

125 <213> ORGANISM: Homo sapiens

127 <400> SEQUENCE: 2

129 Met Glu Ser Ser Phe Ser Phe Gly Val Ile Leu Ala Val Leu Ala Ser
 130 1 5 10 15
 132 Leu Ile Ile Ala Thr Asn Thr Leu Val Ala Val Ala Val Leu Leu Leu
 133 20 25 30
 135 Ile His Lys Asn Asp Gly Val Ser Leu Cys Phe Thr Leu Asn Leu Ala
 136 35 40 45
 138 Val Ala Asp Thr Leu Ile Gly Val Ala Ile Ser Gly Leu Leu Thr Asp
 139 50 55 60
 141 Gln Leu Ser Ser Pro Ser Arg Pro Thr Gln Lys Thr Leu Cys Ser Leu
 142 65 70 75 80
 144 Arg Met Ala Phe Val Thr Ser Ser Ala Ala Ser Val Leu Thr Val
 145 85 90 95
 147 Met Leu Ile Thr Phe Asp Arg Tyr Leu Ala Ile Lys Gln Pro Phe Arg
 148 100 105 110
 150 Tyr Leu Lys Ile Met Ser Gly Phe Val Ala Gly Ala Cys Ile Ala Gly
 151 115 120 125

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153 Leu Trp Leu Val Ser Tyr Leu Ile Gly Phe Leu Pro Leu Gly Ile Pro
154      130                      135                      140
156 Met Phe Gln Gln Thr Ala Tyr Lys Gly Gln Cys Ser Phe Phe Ala Val
157 145                      150                      155                      160
159 Phe His Pro His Phe Val Leu Thr Leu Ser Cys Val Gly Phe Phe Pro
160                      165                      170                      175
162 Ala Met Leu Leu Phe Val Phe Phe Tyr Cys Asp Met Leu Lys Ile Ala
163                      180                      185                      190
165 Ser Met His Ser Gln Gln Ile Arg Lys Met Glu His Ala Gly Ala Met
166      195                      200                      205
168 Ala Gly Gly Tyr Arg Ser Pro Arg Thr Pro Ser Asp Phe Lys Ala Leu
169      210                      215                      220
171 Arg Thr Val Ser Val Leu Ile Gly Ser Phe Ala Leu Ser Trp Thr Pro
172 225                      230                      235                      240
174 Phe Leu Ile Thr Gly Ile Val Gln Val Ala Cys Gln Glu Cys His Leu
175                      245                      250                      255
177 Tyr Leu Val Leu Glu Arg Tyr Leu Trp Leu Leu Gly Val Gly Asn Ser
178                      260                      265                      270
180 Leu Leu Asn Pro Leu Ile Tyr Ala Tyr Trp Gln Lys Glu Val Arg Leu
181                      275                      280                      285
183 Gln Leu Tyr His Met Ala Leu Gly Val Lys Lys Val Leu Thr Ser Phe
184      290                      295                      300
186 Leu Leu Phe Leu Ser Ala Arg Asn Cys Gly Pro Glu Arg Pro Arg Glu
187 305                      310                      315                      320
189 Ser Ser Cys His Ile Val Thr Ile Ser Ser Ser Glu Phe Asp Gly
190                      325                      330                      335

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192 <210> SEQ ID NO: 3

193 <211> LENGTH: 657

194 <212> TYPE: DNA

195 <213> ORGANISM: Homo sapiens

197 <400> SEQUENCE: 3

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198 cagcgcgagc gccttcattg tgaagggtgc catgcgctgg cagtgtctgc gtgccaccgc      60
200 gtgcacctgg agcgaggtag ggcagagcac cgccagcggc agcacgaagc ccacggcatg      120
202 gagcgtggcg gtgaaggctg cgaagcgcgg acgctcaggc tcgggcgga ggcgcagcga      180
204 acaggacgag aaggcgctgc ttagaccaag ccacgagcag ccaagtgcag cgctgagaa      240
206 ggccagcgac tgtccccagg cacagcccag cagcaggccg gcatagcgcg gtcgcaggcg      300
208 tccggcgtag cgcagtggga agcccactgc cagccactgg tctgcgctca gcgccgccac      360
210 gctcagcgcc gcgttgagcg ccaggaaggt gtccaggaag ccaatgactt ggcattgcgc      420
212 gggcgccgac ggtgtccgcc cgcgcacac accgagcagc gtgaagggca tgtccagcgc      480
214 cgccagcagc aggtggccca gagacagatt caccaggagg acgctgagg ctcgagtgcg      540
216 gagctcagcg ctgtaggcgc aacaaagcag caccagtgcg ttgatagca gcgccacggc      600
218 cagtaccatc accaggagac ccgccagcag cgcctcgccg gggcccatgg cgctagc      657

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221 <210> SEQ ID NO: 4

222 <211> LENGTH: 217

223 <212> TYPE: PRT

224 <213> ORGANISM: Homo sapiens

226 <400> SEQUENCE: 4

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228 Ser Ala Met Gly Pro Gly Glu Ala Leu Leu Ala Gly Leu Leu Val Met
229 1                      5                      10                      15

```

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DATE: 10/02/2001

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Input Set : A:\00431PHRM293.ST25.txt

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231 Val Leu Ala Val Ala Leu Leu Ser Asn Ala Leu Val Leu Leu Cys Cys
232           20           25           30
234 Ala Tyr Ser Ala Glu Leu Arg Thr Arg Ala Ser Gly Val Leu Leu Val
235           35           40           45
237 Asn Leu Ser Leu Gly His Leu Leu Leu Ala Ala Leu Asp Met Pro Phe
238           50           55           60
240 Thr Leu Leu Gly Val Met Arg Gly Arg Thr Pro Ser Ala Pro Gly Ala
241 65           70           75           80
243 Cys Gln Val Ile Gly Phe Leu Asp Thr Phe Leu Ala Ser Asn Ala Ala
244           85           90           95
246 Leu Ser Val Ala Ala Leu Ser Ala Asp Gln Trp Leu Ala Val Gly Phe
247           100          105          110
249 Pro Leu Arg Tyr Ala Gly Arg Leu Arg Pro Arg Tyr Ala Gly Leu Leu
250           115          120          125
252 Leu Gly Cys Ala Trp Gly Gln Ser Leu Ala Phe Ser Gly Ala Ala Leu
253           130          135          140
255 Gly Cys Ser Trp Leu Gly Tyr Ser Ser Ala Phe Ala Ser Cys Ser Leu
256 145          150          155          160
258 Arg Leu Pro Pro Glu Pro Glu Arg Pro Arg Phe Ala Ala Phe Thr Ala
259           165          170          175
261 Thr Leu His Ala Val Gly Phe Val Leu Pro Leu Ala Val Leu Cys Leu
262           180          185          190
264 Thr Ser Leu Gln Val His Arg Val Ala Arg Arg His Cys Gln Arg Met
265           195          200          205
267 Asp Thr Val Thr Met Lys Ala Leu Ala
268           210          215

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270 <210> SEQ ID NO: 5

271 <211> LENGTH: 222

272 <212> TYPE: DNA

273 <213> ORGANISM: Homo sapiens

275 <400> SEQUENCE: 5

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276 tgtgcagggtg tgatctccat tcctttgtac atcoctcaca cgctgttcga atgggatttt      60
278 ggaaaggaaa tctgtgtatt ttggctcact actgactatc tgttatgtac agcatctgta      120
280 tataacattg tctcatcag ctatgatcga tacctgtcag tctcaaatgc tgtaagtcga      180
282 acacattaat ttatccccct tagaagatta tgtaaagtga ta                          222

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285 <210> SEQ ID NO: 6

286 <211> LENGTH: 73

287 <212> TYPE: PRT

288 <213> ORGANISM: Homo sapiens

290 <400> SEQUENCE: 6

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292 Cys Ala Gly Val Ile Ser Ile Pro Leu Tyr Ile Pro His Thr Leu Phe
293 1           5           10          15
295 Glu Trp Asp Phe Gly Lys Glu Ile Cys Val Phe Trp Leu Thr Thr Asp
296           20           25           30
298 Tyr Leu Leu Cys Thr Ala Ser Val Tyr Asn Ile Val Leu Ile Ser Tyr
299           35           40           45
301 Asp Arg Tyr Leu Ser Val Ser Asn Ala Val Ser Arg Thr His Phe Ile
302           50           55           60
304 Pro Leu Arg Arg Leu Cys Lys Cys Ile

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TIME: 09:23:49

Input Set : A:\00431PHRM293.ST25.txt

Output Set: N:\CRF3\10022001\I714449A.raw

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307 <210> SEQ ID NO: 7
308 <211> LENGTH: 507
309 <212> TYPE: DNA
310 <213> ORGANISM: Homo sapiens
312 <400> SEQUENCE: 7
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315 cagcggggac agggcgggtca ggagcagcag ccagggtccct gcacacgcgg ccaccgcgta      120
317 acgacggcgg cgccagcgct tggagctgag cgggtacagg atccccagga agcgctccac      180
319 gctgatacag gtcattggtga ggatgctgga atacatgttt gcgtaaaagg ccacgggtcac      240
321 cacgttgcaa agcagcacc cgaataccca gtggtggcgg ttgcaatggt agtagatttg      300
323 gaaaggcaac acgctggcca gcatcaggtc cgtgacgctc aggttgatca tgaagatgac      360
325 cgacggggat ctggggcccca tgcgcgggca cagcaccacac agagagaaga ggttgcccgg      420
327 gatgctgacc gccgccacca gcgagtacac cacgggcagg gccaccgcga tcgcccgggt      480
329 ccgcagcatc tgcagcgctc cgttgctc
332 <210> SEQ ID NO: 8
333 <211> LENGTH: 169
334 <212> TYPE: PRT
335 <213> ORGANISM: Homo sapiens
337 <400> SEQUENCE: 8
339 Asp Asn Ala Thr Leu Gln Met Leu Arg Asn Pro Ala Ile Ala Val Ala
340 1          5          10          15
342 Leu Pro Val Val Tyr Ser Leu Val Ala Ala Val Ser Ile Pro Gly Asn
343          20          25          30
345 Leu Phe Ser Leu Trp Val Leu Cys Arg Arg Met Gly Pro Arg Ser Pro
346          35          40          45
348 Ser Val Ile Phe Met Ile Asn Leu Ser Val Thr Asp Leu Met Leu Ala
349          50          55          60
351 Ser Val Leu Pro Phe Gln Ile Tyr Tyr His Cys Asn Arg His His Trp
352 65          70          75          80
354 Val Phe Gly Val Leu Cys Asn Leu Val Val Thr Val Ala Phe Tyr Ala
355          85          90          95
357 Asn Met Tyr Ser Ser Ile Leu Thr Met Thr Cys Ile Ser Val Glu Arg
358          100         105         110
360 Phe Leu Gly Ile Leu Tyr Pro Leu Ser Ser Lys Arg Trp Arg Arg Arg
361          115         120         125
363 Arg Tyr Ala Val Ala Ala Cys Ala Gly Thr Trp Leu Leu Leu Leu Thr
364          130         135         140
366 Ala Leu Ser Pro Leu Ala Arg Thr Asp Leu Thr Tyr Pro Val His Ala
367 145         150         155         160
369 Leu Gly Ile Ile Thr Cys Phe Asp Val
370          165
372 <210> SEQ ID NO: 9
373 <211> LENGTH: 270
374 <212> TYPE: DNA
375 <213> ORGANISM: Homo sapiens
377 <400> SEQUENCE: 9
378 cccatgttcc tgctcctggg cagcctcacy ttgtcggatc tgctggcagg cgccgcctac      60
380 gccgcccaaca tctactgtc ggggcccgtc acgctgaaac tgtccccgc gctctggttc      120

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Use of n and / or Xaa has been detected in the
Sequence Listing. Review the Sequence Listing
to ensure a corresponding explanation is present
in the <220> to <223> fields of each sequence
using n or Xaa.

VERIFICATION SUMMARY

DATE: 10/02/2001

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Input Set : A:\00431PHRM293.ST25.txt

Output Set: N:\CRF3\10022001\I714449A.raw

L:612 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:624 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:651 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:654 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:672 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:675 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:985 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:1029 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1032 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:2026 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58